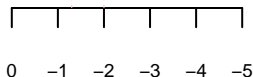


Oecy.Fem-Mal_difpos domain

| | |
|-----------------|------------------|
| Ala_racemase_N | p = 0.017 n = 1 |
| Arginase | p = 0.017 n = 1 |
| ATP-cone | p = 0.017 n = 1 |
| CAF1A | p = 0.017 n = 1 |
| dUTPase | p = 0.017 n = 1 |
| eIF_4EBP | p = 0.017 n = 1 |
| Glyco_hydro_2 | p = 0.017 n = 1 |
| Glyco_hydro_2_C | p = 0.017 n = 1 |
| Glyco_hydro_2_N | p = 0.017 n = 1 |
| Glyoxalase_4 | p = 0.017 n = 1 |
| HgmA | p = 0.017 n = 1 |
| LRR_8 | p = 0.017 n = 5 |
| Mak16 | p = 0.017 n = 1 |
| MCM | p = 0.017 n = 2 |
| MCM_N | p = 0.017 n = 2 |
| MCM_OB | p = 0.017 n = 2 |
| Methyltransf_4 | p = 0.017 n = 1 |
| Microtub_bind | p = 0.017 n = 1 |
| NOC3p | p = 0.017 n = 1 |
| NPR2 | p = 0.017 n = 1 |
| Phos_pyr_kin | p = 0.017 n = 1 |
| Ribonuc_red_lgC | p = 0.017 n = 1 |
| Ribonuc_red_lgN | p = 0.017 n = 1 |
| Ribosomal_L20 | p = 0.017 n = 1 |
| SSF | p = 0.017 n = 2 |
| TIR | p = 0.017 n = 2 |
| Tma16 | p = 0.017 n = 1 |
| Cdc6_C | p = 0.0218 n = 1 |
| CP2 | p = 0.0218 n = 1 |
| DNA_ligase_A_C | p = 0.0218 n = 1 |



log(p)

n = 72 genes | p adjust BH TRUE